



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/764,415
Source: IFWO
Date Processed by STIC: 7/6/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

~~TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER~~
~~VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND~~
~~TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:~~

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box-1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/764,415

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☐ **Wrapped Nucleics
Wrapped Aminos** The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 ☐ **Invalid Line Length** The rules require that a line not exceed 72 characters in length. This includes white spaces.

- 3 ☐ **Misaligned Amino
Numbering** The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters instead.

- 4 ☐ **Non-ASCII** The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

- 5 ☒ **Variable Length** Sequence(s) 4 contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 ☐ **PatentIn 2.0
"bug"** A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

- 7 ☐ **Skipped Sequences
(OLD RULES)** Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

- 8 ☐ **Skipped Sequences
(NEW RULES)** Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

- 9 ☐ **Use of n's or Xaa's
(NEW RULES)** Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

- 10 ☐ **Invalid <213>
Response** Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

- 11 ☐ **Use of <220>** Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

- 12 ☐ **PatentIn 2.0
"bug"** Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 ☐ **Misuse of n/Xaa** "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFWO

RAW SEQUENCE LISTING

DATE: 07/06/2004

PATENT APPLICATION: US/10/764,415

TIME: 10:10:35

Input Set : A:\P11073us seq list.ST25.txt

Output Set: N:\CRF4\07062004\J764415.raw

5 <110> APPLICANT: Lorantis Ltd.
 9 <120> TITLE OF INVENTION: Modulations of Notch signalling for use in Immunotherapy
 13 <130> FILE REFERENCE: P011073US
 17 <140> CURRENT APPLICATION NUMBER: 10/764,415
 C--> 19 <141> CURRENT FILING DATE: 2004-01-23
 23 <150> PRIOR APPLICATION NUMBER: GB0118153.6
 25 <151> PRIOR FILING DATE: 2001-07-01
 29 <150> PRIOR APPLICATION NUMBER: GB0207930.9
 31 <151> PRIOR FILING DATE: 2002-04-05
 35 <150> PRIOR APPLICATION NUMBER: GB0212283.6
 37 <151> PRIOR FILING DATE: 2002-05-28
 41 <150> PRIOR APPLICATION NUMBER: GB0212282.8
 43 <151> PRIOR FILING DATE: 2002-05-28
 47 <160> NUMBER OF SEQ ID NOS: 40
 51 <170> SOFTWARE: PatentIn version 3.0
 55 <210> SEQ ID NO: 1
 57 <211> LENGTH: 43
 59 <212> TYPE: PRT
 61 <213> ORGANISM: artificial sequence
 65 <220> FEATURE:
 67 <221> NAME/KEY: MISC_FEATURE
 69 <222> LOCATION: (2)..(9)
 71 <223> OTHER INFORMATION: N is any amino acid residue
 75 <220> FEATURE:
 77 <221> NAME/KEY: MISC_FEATURE
 79 <222> LOCATION: (11)..(13)
 81 <223> OTHER INFORMATION: N is any amino acid residue
 85 <220> FEATURE:
 87 <221> NAME/KEY: MISC_FEATURE
 89 <222> LOCATION: (15)..(25)
 91 <223> OTHER INFORMATION: N is any amino acid residue
 95 <220> FEATURE:
 97 <221> NAME/KEY: MISC_FEATURE
 99 <222> LOCATION: (27)..(33)
 101 <223> OTHER INFORMATION: N is any amino acid residue
 105 <220> FEATURE:
 107 <221> NAME/KEY: MISC_FEATURE
 109 <222> LOCATION: (35)..(42)
 111 <223> OTHER INFORMATION: N is any amino acid residue
 115 <400> SEQUENCE: 1
 W--> 117 Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
 118 1 5 10 15
 120 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa

pp 1,2,4-6
 Does Not Comply
 Corrected Diskette Needed

see p.6 for error explanation

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Input Set : A:\P11073us seq list.ST25.txt

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```

121          20          25          30
123 Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
124          35          40
126 <210> SEQ ID NO: 2
128 <211> LENGTH: 43
130 <212> TYPE: PRT
132 <213> ORGANISM: Artificial sequence
136 <220> FEATURE:
138 <221> NAME/KEY: MISC_FEATURE
140 <222> LOCATION: (2)..(4)
142 <223> OTHER INFORMATION: X is any amino acid residue
146 <220> FEATURE:
148 <221> NAME/KEY: MISC_FEATURE
150 <222> LOCATION: (5)..(6)
152 <223> OTHER INFORMATION: X is any aromatic amino acid residue
156 <220> FEATURE:
158 <221> NAME/KEY: MISC_FEATURE
160 <222> LOCATION: (7)..(9)
162 <223> OTHER INFORMATION: X is any amino acid residue
166 <220> FEATURE:
168 <221> NAME/KEY: MISC_FEATURE
170 <222> LOCATION: (11)..(13)
172 <223> OTHER INFORMATION: X is any amino acid residue
176 <220> FEATURE:
178 <221> NAME/KEY: MISC_FEATURE
180 <222> LOCATION: (15)..(15)
182 <223> OTHER INFORMATION: X is any basic amino acid residue
186 <220> FEATURE:
188 <221> NAME/KEY: MISC_FEATURE
190 <222> LOCATION: (16)..(16)
192 <223> OTHER INFORMATION: X is any non polar amino acid residue
196 <220> FEATURE:
198 <221> NAME/KEY: MISC_FEATURE
200 <222> LOCATION: (17)..(17)
202 <223> OTHER INFORMATION: X is any basic amino acid residue
206 <220> FEATURE:
208 <221> NAME/KEY: MISC_FEATURE
210 <222> LOCATION: (18)..(19)
212 <223> OTHER INFORMATION: X is any (acid or amide) amino acid residue
216 <220> FEATURE:
218 <221> NAME/KEY: MISC_FEATURE
220 <222> LOCATION: (20)..(20)
222 <223> OTHER INFORMATION: X is any amino acid residue
226 <220> FEATURE:
228 <221> NAME/KEY: MISC_FEATURE
230 <222> LOCATION: (21)..(21)
232 <223> OTHER INFORMATION: X is any aromatic amino acid residue
236 <220> FEATURE:
238 <221> NAME/KEY: MISC_FEATURE

```

see p. 6

? what is this? Xaa can only represent a single amino acid.

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Input Set : A:\P11073us seq list.ST25.txt

Output Set: N:\CRF4\07062004\J764415.raw

240 <222> LOCATION: (22)..(22)
 242 <223> OTHER INFORMATION: X is any non polar amino acid residue
 246 <220> FEATURE:
 248 <221> NAME/KEY: MISC_FEATURE
 250 <222> LOCATION: (23)..(23)
 252 <223> OTHER INFORMATION: X is any aromatic amino acid residue
 256 <220> FEATURE:
 258 <221> NAME/KEY: MISC_FEATURE
 260 <222> LOCATION: (24)..(25)
 262 <223> OTHER INFORMATION: X is any amino acid residue
 266 <220> FEATURE:
 268 <221> NAME/KEY: MISC_FEATURE
 270 <222> LOCATION: (27)..(29)
 272 <223> OTHER INFORMATION: X is any amino acid residue
 276 <220> FEATURE:
 278 <221> NAME/KEY: MISC_FEATURE
 280 <222> LOCATION: (30)..(30)
 282 <223> OTHER INFORMATION: X is any non polar amino acid residue
 286 <220> FEATURE:
 288 <221> NAME/KEY: MISC_FEATURE
 290 <222> LOCATION: (31)..(33)
 292 <223> OTHER INFORMATION: X is any amino acid residue
 296 <220> FEATURE:
 298 <221> NAME/KEY: MISC_FEATURE
 300 <222> LOCATION: (35)..(36)
 302 <223> OTHER INFORMATION: X is any amino acid residue
 306 <220> FEATURE:
 308 <221> NAME/KEY: MISC_FEATURE
 310 <222> LOCATION: (38)..(38)
 312 <223> OTHER INFORMATION: X is any aromatic amino acid residue
 316 <220> FEATURE:
 318 <221> NAME/KEY: MISC_FEATURE
 320 <222> LOCATION: (37)..(37)
 322 <223> OTHER INFORMATION: X is any non polar amino acid residue
 326 <220> FEATURE:
 328 <221> NAME/KEY: MISC_FEATURE
 330 <222> LOCATION: (39)..(39)
 332 <223> OTHER INFORMATION: X is any amino acid residue
 336 <220> FEATURE:
 338 <221> NAME/KEY: MISC_FEATURE
 340 <222> LOCATION: (40)..(40)
 342 <223> OTHER INFORMATION: X is any non polar amino acid residue
 346 <220> FEATURE:
 348 <221> NAME/KEY: MISC_FEATURE
 350 <222> LOCATION: (41)..(42)
 352 <223> OTHER INFORMATION: X is any amino acid residue
 356 <400> SEQUENCE: 2

W--> 358 Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa

359 1

5

10

15

RAW SEQUENCE LISTING

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Input Set : A:\P11073us seq list.ST25.txt

Output Set : N:\CRF4\07062004\J764415.raw

361 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa
362 20 25 30
364 Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
365 35 40
367 <210> SEQ ID NO: 3
369 <211> LENGTH: 43
371 <212> TYPE: PRT
373 <213> ORGANISM: artificial sequence *see p. 6*
377 <220> FEATURE:
379 <221> NAME/KEY: misc_feature
381 <222> LOCATION: (2)..(4) ✓
383 <223> OTHER INFORMATION: X is any amino acid residue
387 <220> FEATURE:
389 <221> NAME/KEY: misc_feature
391 <222> LOCATION: (7)..(9) ✓
393 <223> OTHER INFORMATION: X is any amino acid residue
397 <220> FEATURE:
399 <221> NAME/KEY: misc_feature
401 <222> LOCATION: (11)..(13) ✓
403 <223> OTHER INFORMATION: X is any amino acid residue
407 <220> FEATURE:
409 <221> NAME/KEY: misc_feature
411 <222> LOCATION: (18)..(18) ✓
413 <223> OTHER INFORMATION: X is either aspartic acid *or?* of asparagine
417 <220> FEATURE:
419 <221> NAME/KEY: misc_feature
421 <222> LOCATION: (20)..(20) ✓
423 <223> OTHER INFORMATION: X is any amino acid residue
427 <220> FEATURE:
429 <221> NAME/KEY: misc_feature
431 <222> LOCATION: (24)..(25) ✓
433 <223> OTHER INFORMATION: X is any amino acid residue
437 <220> FEATURE:
439 <221> NAME/KEY: misc_feature
441 <222> LOCATION: (27)..(29) ✓
443 <223> OTHER INFORMATION: X is any amino acid residue
447 <220> FEATURE:
449 <221> NAME/KEY: misc_feature
451 <222> LOCATION: (31)..(33) ✓
453 <223> OTHER INFORMATION: X is any amino acid residue
457 <220> FEATURE:
459 <221> NAME/KEY: misc_feature
461 <222> LOCATION: (35)..(36) ✓
463 <223> OTHER INFORMATION: X is any amino acid residue
467 <220> FEATURE:
469 <221> NAME/KEY: misc_feature
471 <222> LOCATION: (39)..(39) ✓
473 <223> OTHER INFORMATION: X is any amino acid residue
477 <220> FEATURE:

RAW SEQUENCE LISTING

DATE: 07/06/2004

PATENT APPLICATION: US/10/764,415

TIME: 10:10:35

Input Set : A:\P11073us seq list.ST25.txt

Output Set: N:\CRF4\07062004\J764415.raw

479 <221> NAME/KEY: misc_feature
 481 <222> LOCATION: (40)..(42)
 483 <223> OTHER INFORMATION: X is any amino acid residue
 487 <400> SEQUENCE: 3
 W--> 489 Cys Xaa Xaa Xaa Tyr Tyr Xaa Xaa Xaa Cys Xaa Xaa Xaa Cys Arg Pro
 490 1 5 10 15
 492 Arg Xaa Asp Xaa Phe Gly His Xaa Xaa Cys Xaa Xaa Xaa Gly Xaa Xaa
 493 20 25 30
 495 Xaa Cys Xaa Xaa Gly Trp Xaa Gly Xaa Xaa Cys
 496 35 40
 498 <210> SEQ ID NO: 4
 500 <211> LENGTH: 18
 502 <212> TYPE: PRT
 504 <213> ORGANISM: artificial sequence
 508 <220> FEATURE:
 510 <221> NAME/KEY: MISC_FEATURE
 512 <222> LOCATION: (1)..(1)
 514 <223> OTHER INFORMATION: X is 1 to 4 amino acids
 518 <220> FEATURE:
 520 <221> NAME/KEY: MISC_FEATURE
 522 <222> LOCATION: (3)..(3)
 524 <223> OTHER INFORMATION: X is 0 to 48 amino acids
 528 <220> FEATURE:
 530 <221> NAME/KEY: MISC_FEATURE
 532 <222> LOCATION: (5)..(5)
 534 <223> OTHER INFORMATION: X is 3 to 12 amino acids
 538 <220> FEATURE:
 540 <221> NAME/KEY: MISC_FEATURE
 542 <222> LOCATION: (7)..(7)
 544 <223> OTHER INFORMATION: X is 1 to 70 amino acids
 548 <220> FEATURE:
 550 <221> NAME/KEY: MISC_FEATURE
 552 <222> LOCATION: (9)..(9)
 554 <223> OTHER INFORMATION: X is 1 to 6 amino acids
 558 <220> FEATURE:
 560 <221> NAME/KEY: MISC_FEATURE
 562 <222> LOCATION: (11)..(11)
 564 <223> OTHER INFORMATION: X is 2 amino acids
 568 <220> FEATURE:
 570 <221> NAME/KEY: MISC_FEATURE
 572 <222> LOCATION: (14)..(14)
 574 <223> OTHER INFORMATION: X is 0 to 21 amino acids
 578 <220> FEATURE:
 580 <221> NAME/KEY: MISC_FEATURE
 582 <222> LOCATION: (16)..(16)
 584 <223> OTHER INFORMATION: X is 2 amino acids
 588 <220> FEATURE:
 590 <221> NAME/KEY: MISC_FEATURE
 592 <222> LOCATION: (18)..(18)

see p.6

*variable length is not permitted
 (see item 5 on
 Error Summary sheet)*

RAW SEQUENCE LISTING ERROR SUMMARY
 PATENT APPLICATION: US/10/764,415

DATE: 07/06/2004
 TIME: 10:10:36

Input Set : A:\P11073us seq list.ST25.txt
 Output Set: N:\CRF4\07062004\J764415.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. ~~2,3,4,5,6,7,8,9,11,12,13,15,16,17,18,19,20,21,22,23,24,25~~
 Seq#:1; Xaa Pos. ~~27,28,29,30,31,32,33,35,36,37,38,39,40,41,42~~
 Seq#:2; Xaa Pos. ~~2,3,4,5,6,7,8,9,11,12,13,15,16,17,18,19,20,21,22,23,24,25~~
 Seq#:2; Xaa Pos. ~~27,28,29,30,31,32,33,35,36,37,38,39,40,41,42~~
 Seq#:3; Xaa Pos. ~~2,3,4,7,8,9,11,12,13,18,20,24,25,27,28,29,31,32,33,35,36~~
 Seq#:3; Xaa Pos. ~~39,41,42~~
 Seq#:4; Xaa Pos. ~~1,3,5,7,9,11,14,16,18~~
 Seq#:39; Xaa Pos. 891

Use of <220> Feature(NEW RULES): *error explanation*

Sequence(s) are missing the <220> Feature and associated headings.

Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:5,6,7,8,9,10,11,12,13,14,15,16,17,*1,2,3,4*

VERIFICATION SUMMARY

DATE: 07/06/2004

PATENT APPLICATION: US/10/764,415

TIME: 10:10:36

Input Set : A:\P11073us seq list.ST25.txt

Output Set: N:\CRF4\07062004\J764415.raw

L:19 M:271 C: Current Filing Date differs, Replaced Current Filing Date
 L:117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
 M:341 Repeated in SeqNo=1
 L:358 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
 M:341 Repeated in SeqNo=2
 L:489 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
 M:341 Repeated in SeqNo=3
 L:600 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
 M:341 Repeated in SeqNo=4
 L:616 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:5, <213>
 ORGANISM:Artificial sequence
 L:616 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:5, <213>
 ORGANISM:Artificial sequence
 L:616 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:5,Line#:616
 L:630 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:6, <213>
 ORGANISM:Artificial sequence
 L:630 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:6, <213>
 ORGANISM:Artificial sequence
 L:630 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:6,Line#:630
 L:644 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:7, <213>
 ORGANISM:Artificial sequence
 L:644 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:7, <213>
 ORGANISM:Artificial sequence
 L:644 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:7,Line#:644
 L:658 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:8, <213>
 ORGANISM:Artificial sequence
 L:658 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:8, <213>
 ORGANISM:Artificial sequence
 L:658 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:8,Line#:658
 L:672 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:9, <213>
 ORGANISM:Artificial sequence
 L:672 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:9, <213>
 ORGANISM:Artificial sequence
 L:672 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:9,Line#:672
 L:686 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:10, <213>
 ORGANISM:Artificial sequence
 L:686 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:10, <213>
 ORGANISM:Artificial sequence
 L:686 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:10,Line#:686
 L:700 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:11, <213>
 ORGANISM:Artificial sequence
 L:700 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:11, <213>
 ORGANISM:Artificial sequence
 L:700 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:11,Line#:700
 L:714 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:12, <213>
 ORGANISM:Artificial sequence
 L:714 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:12, <213>
 ORGANISM:Artificial sequence
 L:714 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:12,Line#:714
 L:728 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:13, <213>
 ORGANISM:Artificial sequence
 L:728 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:13, <213>
 ORGANISM:Artificial sequence

L:728 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:13,Line#:728
L:742 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial sequence
L:742 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial sequence
L:742 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:742
L:756 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:15, <213>
ORGANISM:Artificial Sequence
L:756 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:15, <213>
ORGANISM:Artificial Sequence
L:756 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:15,Line#:756
L:770 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:16, <213>
ORGANISM:Artificial sequence
L:770 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:16, <213>
ORGANISM:Artificial sequence
L:770 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:16,Line#:770
L:784 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:17, <213>
ORGANISM:Artificial sequence
L:784 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:17, <213>
ORGANISM:Artificial sequence.
L:784 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:17,Line#:784

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/764,415

DATE: 07/06/2004

TIME: 10:10:36

Input Set : A:\P11073us seq list.ST25.txt

Output Set: N:\CRF4\07062004\J764415.raw

L:2419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:880